

Query Match 100.0%; Score 1831; DB 17; Length 352;  
Best Local Similarity 100.0%; Pred. No. 1,66-162;  
Matches 352; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MESGGRPSLCOFILGTTSVTTAALYSVYRQKARVSGELGAKKVLHGEDLSKLSLSEAPG 60  
Db 1 MESGGRPSLCOFILGTTSVTTAALYSVYRQKARVSGELGAKKVLHGEDLSKLSLSEAPG 60

Qy 61 KCVPAVIEGAVRSVKETLNSQFVENCCKVIOQLTLOEHKVMNRTHLMNDSKITHOR 120  
Db 61 KCVPAVIEGAVRSVKETLNSQFVENCCKVIOQLTLOEHKVMNRTHLMNDSKITHOR 120

Qy 121 TMTVPFDLVPHEDGDVAVRVLKPLDSVDLGLFTVYEKHPISQSFVDVIGHYISGERPK 180  
Db 121 TMTVPFDLVPHEDGDVAVRVLKPLDSVDLGLFTVYEKHPISQSFVDVIGHYISGERPK 180

Qy 181 GIQETEBMLKVGATLTGVELVLDNNSVRLQPPKQMOYILSSQDFSLQROESSVRLW 240  
Db 181 GIQETEBMLKVGATLTGVELVLDNNSVRLQPPKQMOYILSSQDFSLQROESSVRLW 240

Qy 241 KYLALVFGFATCATLFTILRKQYLOQROERLRLKQMOEFQHEHAQLLSRAKPEDRESLKS 300  
Db 241 KYLALVFGFATCATLFTILRKQYLOQROERLRLKQMOEFQHEHAQLLSRAKPEDRESLKS 300

Qy 301 ACVCLSSFKSCVFLGCHVCSCTECYRALPEPKKCPICROAITRVIPLYNS 352  
Db 301 ACVCLSSFKSCVFLGCHVCSCTECYRALPEPKKCPICROAITRVIPLYNS 352

RESULT 9  
US-09-764-864-801  
; Sequence 801, Application US/09764864  
; Patent No. US20020132753A1  
; GENERAL INFORMATION:  
; APPLICANT: Rosen et al.  
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies  
; FILE REFERENCE: PT223  
; CURRENT APPLICATION NUMBER: US/09/764,864  
; CURRENT FILING DATE: 2001-01-17  
; Prior application data removed - consult PALM or file wrapper  
; NUMBER OF SEQ ID NOS: 1792  
; SOFTWARE: Patentin Ver. 2.0  
; SEQ ID NO 801  
; LENGTH: 392  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
; FEATURE:  
; NAME/KEY: SITE  
; LOCATION: (238)  
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids  
US-09-764-864-801

Query Match 99.7%; Score 1826; DB 9; Length 392;  
Best Local Similarity 99.7%; Pred. No. 5,5e-162;  
Matches 351; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MESGGRPSLCOFILGTTSVTTAALYSVYRQKARVSGELGAKKVLHGEDLSKLSLSEAPG 60  
Db 41 MESGGRPSLCOFILGTTSVTTAALYSVYRQKARVSGELGAKKVLHGEDLSKLSLSEAPG 100

Qy 61 KCVPAVIEGAVRSVKETLNSQFVENCCKVIOQLTLOEHKVMNRTHLMNDSKITHOR 120  
Db 101 KCVPAVIEGAVRSVKETLNSQFVENCCKVIOQLTLOEHKVMNRTHLMNDSKITHOR 160

Qy 121 TMTVPFDLVPHEDGDVAVRVLKPLDSVDLGLFTVYEKHPISQSFVDVIGHYISGERPK 180  
Db 161 TMTVPFDLVPHEDGDVAVRVLKPLDSVDLGLFTVYEKHPISQSFVDVIGHYISGERPK 220

Qy 181 GIQETEBMLKVGATLTGVELVLDNNSVRLQPPKQMOYILSSQDFSLQROESSVRLW 240  
Db 221 GIQETEBMLKVGATLTGVELVLDNNSVRLQPPKQMOYILSSQDFSLQROESSVRLW 280

Qy 241 KYLALVFGFATCATLFTILRKQYLOQROERLRLKQMOEFQHEHAQLLSRAKPEDRESLKS 300

Db 281 KYLALVFGFATCATLFTILRKQYLOQROERLRLKQMOEFQHEHAQLLSRAKPEDRESLKS 340  
Qy 301 ACVCLSSFKSCVFLGCHVCSCTECYRALPEPKKCPICROAITRVIPLYNS 352  
Db 341 ACVCLSSFKSCVFLGCHVCSCTECYRALPEPKKCPICROAITRVIPLYNS 392

RESULT 10  
US-10-024-298A-73  
; Sequence 73, Application US/10024298A  
; Publication No. US20030143540A1  
; GENERAL INFORMATION:  
; APPLICANT: ASAHU KASEI KABUSHIKI KAISHA  
; APPLICANT: AKIO MATSUDA  
; APPLICANT: GOICHI HONDA  
; APPLICANT: SHUJI MURAYAMATSU  
; APPLICANT: YUKIKO NAGANO  
; TITLE OF INVENTION: NF-K B Activating Gene  
; FILE REFERENCE: 1254-0191P  
; CURRENT APPLICATION NUMBER: US/10/024,298A  
; PRIOR FILING DATE: 2003-04-08  
; PRIOR APPLICATION NUMBER: 60/314,385  
; PRIOR FILING DATE: 2001-08-24  
; PRIOR APPLICATION NUMBER: 60/278,641  
; PRIOR FILING DATE: 2001-03-26  
; PRIOR APPLICATION NUMBER: 60/258,315  
; PRIOR FILING DATE: 2000-12-28  
; PRIOR APPLICATION NUMBER: JP254016/2001  
; PRIOR FILING DATE: 2001-08-24  
; PRIOR APPLICATION NUMBER: JP088912/2001  
; PRIOR FILING DATE: 2001-03-26  
; PRIOR APPLICATION NUMBER: JP402288/2000  
; PRIOR FILING DATE: 2000-12-28  
; NUMBER OF SEQ ID NOS: 182  
; SOFTWARE: Patentin Ver. 2.0  
; SEQ ID NO 73  
; LENGTH: 352  
; TYPE: PRT  
; ORGANISM: Homo sapiens  
US-10-024-298A-73

Query Match 99.6%; Score 1824; DB 14; Length 352;  
Best Local Similarity 99.7%; Pred. No. 7,3e-162;  
Matches 351; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MESGGRPSLCOFILGTTSVTTAALYSVYRQKARVSGELGAKKVLHGEDLSKLSLSEAPG 60  
Db 1 MESGGRPSLCOFILGTTSVTTAALYSVYRQKARVSGELGAKKVLHGEDLSKLSLSEAPG 60

Qy 61 KCVPAVIEGAVRSVKETLNSQFVENCCKVIOQLTLOEHKVMNRTHLMNDSKITHOR 120  
Db 61 KCVPAVIEGAVRSVKETLNSQFVENCCKVIOQLTLOEHKVMNRTHLMNDSKITHOR 120

Qy 121 TMTVPFDLVPHEDGDVAVRVLKPLDSVDLGLFTVYEKHPISQSFVDVIGHYISGERPK 180  
Db 121 TMTVPFDLVPHEDGDVAVRVLKPLDSVDLGLFTVYEKHPISQSFVDVIGHYISGERPK 180

Qy 181 GIQETEBMLKVGATLTGVELVLDNNSVRLQPPKQMOYILSSQDFSLQROESSVRLW 240  
Db 181 GIQETEBMLKVGATLTGVELVLDNNSVRLQPPKQMOYILSSQDFSLQROESSVRLW 240

Qy 241 KYLALVFGFATCATLFTILRKQYLOQROERLRLKQMOEFQHEHAQLLSRAKPEDRESLKS 300  
Db 241 KYLALVFGFATCATLFTILRKQYLOQROERLRLKQMOEFQHEHAQLLSRAKPEDRESLKS 300

Qy 301 ACVCLSSFKSCVFLGCHVCSCTECYRALPEPKKCPICROAITRVIPLYNS 352  
Db 301 ACVCLSSFKSCVFLGCHVCSCTECYRALPEPKKCPICROAITRVIPLYNS 352

RESULT 11  
US-10-042-211A-73

CC storing the sequence data on a computer system, and a method for  
CC identifying features of the cDNA sequences using a computer programme.  
CC The cDNAs are useful for expressing secreted proteins or fragments to  
CC obtain antibodies capable of specifically binding to the secreted  
CC proteins. The cDNAs may also be useful in diagnostic, forensic, gene  
CC therapy and chromosome mapping procedures and may be used to design  
CC expression vectors and secretion vectors. The proteins of the invention  
CC may be used to treat diseases including cancer, autoimmune diseases,  
CC cardiovascular disorders, cystic fibrosis, hypothyroidism, immunological  
CC disorders, amyloidosis, brain disorders, skeletal muscle disorders, eye  
CC disorders, obesity, mitochondrial cytopathies, diabetes, atherosclerosis,  
CC neurodegenerative disorders, graft rejection, Alzheimer's disease,  
CC dementia, hyperlipidaemia, septic shock and impotence  
XX  
XX  
SQ Sequence 352 AA;

Query Match 100.0%; Score 1831; DB 3; Length 352;  
Best Local Similarity 100.0%; Pred. No. 1.2e-168; Indels 0; Gaps 0;  
Matches 352; Conservative 0; Mismatches 0;

QY 1 MESSGRPSLCQFILGTTSVTAAALYSYVRQKARVSGELGAKKVVHGEDLKSILSEAPG 60  
DB 1 MESSGRPSLCQFILGTTSVTAAALYSYVRQKARVSGELGAKKVVHGEDLKSILSEAPG 60  
QY 61 KCVPYAVITGAVRSVKETLNSQFVENCCKVQRLTLOEHKQVNRTHLMDCKSIKHOR 120  
DB 61 KCVPYAVITGAVRSVKETLNSQFVENCCKVQRLTLOEHKQVNRTHLMDCKSIKHOR 120  
QY 121 TMTVPDLVPHEDGVDAVAVLKPLDSVDLGLETVYKFPSPISQFTDVI GHYISGERPK 180  
DB 121 TMTVPDLVPHEDGVDAVAVLKPLDSVDLGLETVYKFPSPISQFTDVI GHYISGERPK 180  
QY 181 GIQETEMLKVGATLTGVEGLVDNNNSVRLQPPKQGMQYLISSQDFSLQROESSVRLW 240  
DB 181 GIQETEMLKVGATLTGVEGLVDNNNSVRLQPPKQGMQYLISSQDFSLQROESSVRLW 240  
QY 241 KVALVFGFATCATLFFILRKQYLOQROELRLKQMOEFQHEAQLSRAKPEDRESLKS 300  
DB 241 KVALVFGFATCATLFFILRKQYLOQROELRLKQMOEFQHEAQLSRAKPEDRESLKS 300  
QY 301 ACVCLSSFKSCVFLBEGHVCSTCECYRALPEPKKPCICROAITRVIPLYNS 352  
DB 301 ACVCLSSFKSCVFLBEGHVCSTCECYRALPEPKKPCICROAITRVIPLYNS 352

RESULT 2  
AAE06602  
ID AAE06602 standard; protein; 352 AA.

AAE06602;  
25-SEP-2001 (first entry)  
Human protein having hydrophobic domain, HPI0649.  
Human: hydrophobic domain; gene therapy; nutritional supplement;  
cell proliferation; immunomodulatory; autoimmune disorder; antimicrobial;  
multiple sclerosis; rheumatoid arthritis; insulin-dependent diabetes;  
haematopoiesis; tissue growth activity; Parkinson's disease; cycostatic;  
Huntington's disease; Alzheimer's disease; chemotactic; chemokinetic;  
haemostatic; thrombolytic; tumour growth inhibitor; anabolic;  
contraceptive; antifertility; antiinflammatory.

Homo sapiens.

MO200149728-A2.

12-JUL-2001.

28-DEC-2000; 2000WO-JP009359.

06-JAN-2000; 2000JP-00000585.

06-JAN-2000; 2000JP-00000588.

11-JAN-2000; 2000JP-00002299.  
03-FEB-2000; 2000JP-00026862.  
03-MAR-2000; 2000JP-00058367.  
(PROT-) PROTEGENE INC.  
(SAGA) SAGAMI CHEM RES CENT.  
Kato S, Kimura T;  
WPI; 2001-418355/44.  
N-PSDB; AAD12597.  
Human proteins with hydrophobic domains and the nucleic acids encoding  
them, useful for preventing diagnosing and treating e.g. cancer,  
Alzheimer's and inflammation.

Claim 1; Page 122; 563pp; English.

The present sequence is human protein with hydrophobic domain, HPI0649.  
The polynucleotide and polypeptide of the invention may be used in the  
prevention, diagnosis and treatment of diseases associated with  
inappropriate polypeptide expression. The polynucleotides may be used to  
produce the polypeptide, by inserting the nucleic acids into a host cell  
and culturing the cell to express the protein. The polynucleotides and  
its complementary sequences may also be used as DNA probes in diagnostic  
assays and also used in gene therapy. The polypeptides may also be used  
as antigens in the production of antibodies and in assays to identify  
modulators of polypeptide expression and activity. The polypeptides and  
nucleic acids may be used as nutritional supplements, to modulate  
cytokine and cell proliferation activity, to modulate immune stimulation  
or suppression (e.g. for the treatment of microbial infections and  
autoimmune disorders such as multiple sclerosis, rheumatoid arthritis and  
insulin-dependent diabetes), to modulate haematopoiesis, to modulate  
tissue growth activity (e.g. for the treatment of Parkinson's disease,  
Huntington's disease and Alzheimer's disease), to modulate activin and  
inhibin activity (e.g. for controlling fertility), to modulate  
chemotactic and chemokinetic activity, to modulate haemostatic and  
thrombolytic activity, to modulate receptor ligand activity, to modulate  
inflammation and to inhibit tumour growth

SO Sequence 352 AA;  
Query Match 100.0%; Score 1831; DB 4; Length 352;  
Best Local Similarity 100.0%; Pred. No. 1.2e-168; Indels 0; Gaps 0;  
Matches 352; Conservative 0; Mismatches 0;

QY 1 MESSGRPSLCQFILGTTSVTAAALYSYVRQKARVSGELGAKKVVHGEDLKSILSEAPG 60  
DB 1 MESSGRPSLCQFILGTTSVTAAALYSYVRQKARVSGELGAKKVVHGEDLKSILSEAPG 60  
QY 61 KCVPYAVITGAVRSVKETLNSQFVENCCKVQRLTLOEHKQVNRTHLMDCKSIKHOR 120  
DB 61 KCVPYAVITGAVRSVKETLNSQFVENCCKVQRLTLOEHKQVNRTHLMDCKSIKHOR 120  
QY 121 TMTVPDLVPHEDGVDAVAVLKPLDSVDLGLETVYKFPSPISQFTDVI GHYISGERPK 180  
DB 121 TMTVPDLVPHEDGVDAVAVLKPLDSVDLGLETVYKFPSPISQFTDVI GHYISGERPK 180  
QY 181 GIQETEMLKVGATLTGVEGLVDNNNSVRLQPPKQGMQYLISSQDFSLQROESSVRLW 240  
DB 181 GIQETEMLKVGATLTGVEGLVDNNNSVRLQPPKQGMQYLISSQDFSLQROESSVRLW 240  
QY 241 KVALVFGFATCATLFFILRKQYLOQROELRLKQMOEFQHEAQLSRAKPEDRESLKS 300  
DB 241 KVALVFGFATCATLFFILRKQYLOQROELRLKQMOEFQHEAQLSRAKPEDRESLKS 300  
QY 301 ACVCLSSFKSCVFLBEGHVCSTCECYRALPEPKKPCICROAITRVIPLYNS 352  
DB 301 ACVCLSSFKSCVFLBEGHVCSTCECYRALPEPKKPCICROAITRVIPLYNS 352

RESULT 3  
ABB50174